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Sequence Listing was accepted.

See attached Validation Report.

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217-9197 (toll free).

Reviewer: markspencer

Timestamp: [year=2009; month=8; day=27; hr=12; min=53; sec=30; ms=213; ]

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Application No: 10578840

Version No: 2.0

Input Set:

Output Set:

Started: 2009-07-30 11:00:11.438

Finished: 2009-07-30 11:00:19.922

Elapsed: 0 hr(s) 0 min(s) 8 sec(s) 484 ms

Total Warnings: 122

Total Errors: 2

No. of SeqIDs Defined: 122

Actual SeqID Count: 122

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
W 213	Artificial or Unknown found in <213> in SEQ ID (12)
W 213	Artificial or Unknown found in <213> in SEQ ID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (14)
W 213	Artificial or Unknown found in <213> in SEQ ID (15)
W 213	Artificial or Unknown found in <213> in SEQ ID (16)
W 213	Artificial or Unknown found in <213> in SEQ ID (17)
W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)

**Input Set:**

**Output Set:**

**Started:** 2009-07-30 11:00:11.438  
**Finished:** 2009-07-30 11:00:19.922  
**Elapsed:** 0 hr(s) 0 min(s) 8 sec(s) 484 ms  
**Total Warnings:** 122  
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**No. of SeqIDs Defined:** 122  
**Actual SeqID Count:** 122

Error code	Error Description
	This error has occurred more than 20 times, will not be displayed
E 257	Invalid sequence data feature in <221> in SEQ ID (122)
E 257	Invalid sequence data feature in <221> in SEQ ID (122)

# SEQUENCE LISTING

<110> KIKUCHI, YASUFUMI  
 UNO, SHINSUKE  
 KINOSHITA, YASUKO  
 IIJIMA, SHIGEYUKI  
 FUKUSHIMA, NAOSHI  
 TSUCHIYA, MASAYUKI

<120> HUMANIZED ANTI-CD47 ANTIBODY

<130> 060641-0113

<140> 10578840

<141> 2009-07-30

<150> PCT/JP2004/016744

<151> 2004-11-11

<150> JP 2003-381406

<151> 2003-11-11

<160> 122

<170> PatentIn version 3.5

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<220>

<223> Description of Artificial Sequence: Synthetic polynucleotide

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<223> Description of Artificial Sequence: Synthetic polynucleotide

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gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag      96
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc ttc      144
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
      15              20              25

gcc aac cat gtt att cac tgg gtg cga cag gcc cct gga caa ggg ctt      192
Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
      30              35              40              45

gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat      240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
      50              55              60

gag aag ttc aag gac aga gtc acg atg acc cgg gac acg tcc acg agc      288
Glu Lys Phe Lys Asp Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser
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aca gtc tac atg gag ttg agc agt ctc aga tct gag gac acg gcc gtc      336
Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
      80              85              90

tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa      384
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
      95              100              105

gga acc ctg gtc acc gtc tcc tca ggtgagtggga tccgcg      424

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Gly Thr Leu Val Thr Val Ser Ser  
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<213> Artificial Sequence

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15 20 25

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Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu  
30 35 40 45

gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240  
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn  
50 55 60

gag aag ttc aag gac aga gtc acg atg acc tca gac acg tcc acg agc 288  
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Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val  
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Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Trp Gly Gln  
95 100 105

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Gly Thr Leu Val Thr Val Ser Ser  
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<210> 11

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<223> Description of Artificial Sequence: Synthetic oligonucleotide

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<210> 12

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<223> Description of Artificial Sequence: Synthetic oligonucleotide

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<211> 424



<212> DNA

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<223> Description of Artificial Sequence: Synthetic plasmid  
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gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag	96
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys	
-1 1 5 10	

cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc ttc	144
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe	
15 20 25	

acc aac cat gtt att cac tgg gtg cga cag gcc cct gga caa ggg ctt	192
Thr Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu	
30 35 40 45	

gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat	240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn	
50 55 60	

gag aag ttc aag gac aga gtc acg atg acc tca gac acg tcc acg agc	288
Glu Lys Phe Lys Asp Arg Val Thr Met Thr Ser Asp Thr Ser Thr Ser	
65 70 75	

aca gtc tac atg gag ttg agc agt ctc aga tct gag gac acg gcc gtc	336
Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val	
80 85 90	

tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa	384
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln	
95 100 105	

gga acc ctg gtc acc gtc tcc tca ggtgagtggg tccgcg	424
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110 115	

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Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe  
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gcc aac cat gtt att cac tgg gtg cga cag gcc cct gga caa ggg ctt 192  
 Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu  
 30 35 40 45

gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240  
 Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn  
 50 55 60

gag aag ttc aag gac aaa gtc acg atg acc tca gac acg tcc acg agc 288  
 Glu Lys Phe Lys Asp Lys Val Thr Met Thr Ser Asp Thr Ser Thr Ser  
 65 70 75

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 Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val  
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tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa 384  
 Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln  
 95 100 105

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<210> 17

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<223> Description of Artificial Sequence: Synthetic oligonucleotide

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<223> Description of Artificial Sequence: Synthetic plasmid  
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Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys	
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Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe	
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gcc aac cat gtt att cac tgg gtg cga cag gcc cct gga caa ggg ctt	192
Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu	
30 35 40 45	

gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat	240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn	
50 55 60	

gag aag ttc aag gac aga gtc acg ctg acc tca gac acg tcc acg agc	288
Glu Lys Phe Lys Asp Arg Val Thr Leu Thr Ser Asp Thr Ser Thr Ser	
65 70 75	

aca gtc tac atg gag ttg agc agt ctc aga tct gag gac acg gcc gtc	336
Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val	
80 85 90	

tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa	384
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln	
95 100 105	

gga acc ctg gtc acc gtc tcc tca ggtgagtgga tccgcg	424
Gly Thr Leu Val Thr Val Ser Ser	
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